Amendments to the Specification:

Please replace the paragraph beginning at page 16, line 31 with the following:

      Figure 16: Grafting Bet v 1 specific surface area I. Amino acid alignment of Dau c 1 and Bet v 1. Position no. above sequences refers to Dau c 1 (T14325) (SEQ ID NO: 4). Positions no. below sequences refers to Bet v 1 (z80104) (SEQ ID NO: 5). Black background shows positions in the polypeptide sequences having identical amino acid residues. Grey background shows positions in the polypeptide sequences having homologous amino acid residues. Amino acid positions that in example 1A are targeted for introduction of secondary mutations are indicated with triple bars on black background and amino acid positions for introduction of primary mutations are indicated with black arrows.

Please replace the paragraph beginning at page 17, line 7 with the following:

      Figure 17: Grafting Bet v 1 specific surface area II. Amino acid alignment of Dau c 1 and Bet v 1. Position no. above sequences refers to Dau c 1 (T14325) (SEQ ID NO: 6). Positions no. below sequences refers to Bet v 1 (z80104) (SEQ ID NO: 5). Black background shows positions in the polypeptide sequences having identical amino acid residues. Grey background shows positions in the polypeptide sequences having homologous amino acid residues. Amino acid positions that in example 1B are targeted for introduction of secondary mutations are indicated with triple bars on black background and amino acid positions for introduction of primary mutations are indicated with black arrows.

Please replace the paragraph beginning at page 17, line 17 with the following:

      Figure 18: Amino acid alignment of Lep d 2 and Der p 2. Position no. above sequences refers to Lep d 2 (S66499) (SEQ ID NO: 7). Positions no. below sequences refers to Der p 2 (P49278) (SEQ ID NO: 8). Black background shows positions in the polypeptide sequences having identical amino acid residues. Grey background shows positions in the polypeptide sequences having homologous amino acid residues. Amino acid positions that in example 2A are targeted for introduction of primary mutations are indicated with white arrows on black background. Amino acid positions that in example 2B are targeted for introduction of primary mutations are indicated with black arrows on white background. Amino acid positions suggested for introduction of secondary mutations in either example 2A or 2B are indicated with triple bars on black background.

Please replace the paragraph beginning at page 18, line 28 with the following:

Figure 21: Amino acid alignment of Mal d 1 (2620) and Bet v 1.2801. Position no. above sequences refers to Mal d 1 (accession no. AJ488060) (SEQ ID NO: 9). Positions no. below sequences refers to Bet v 1 (accession no. Z80104) (SEQ ID NO: 5). Black background shows positions in the polypeptide sequences having identical amino acid residues. Grey background shows positions in the polypeptide sequences having homologous amino acid residues. Amino acid positions that in example 3 are targeted for introduction of secondary mutations are indicated with triple bars on black background and amino acid positions for introduction of primary mutations are indicated with black arrows.

Please replace the paragraph beginning at page 19, line 4 with the following:

Figure 22: Amino acid alignment of Gly d 2 and Der p 2. Position no. above sequences refers to Gly d 2 (AJ272216) (SEQ ID NO: 10). Positions no. below sequences refers to Der p 2 (P49278) (SEQ ID NO: 8). Black background shows positions in the polypeptide sequences having identical amino acid residues. Grey background shows positions in the polypeptide sequences having homologous amino acid residues. Amino acid positions that in example 4 are targeted for introduction of secondary mutations are indicated with triple bars on black background and amino acid positions for introduction of primary mutations are indicated with black arrows.

After page 65 and before the claims, please insert a paper copy of the Sequence Listing.